1631



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/923,444A

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SEQUENCE LISTING

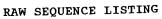
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4 (1) GENERAL INFORMATION:
            (i) APPLICANT: LI, YI
     6
                            FLEISCHMANN, ROBERT
     7
           (ii) TITLE OF INVENTION: NEUROTRANSMITTER TRANSPORTER
     9
          (iii) NUMBER OF SEQUENCES: 6
    11
            (iv) CORRESPONDENCE ADDRESS:
                                                             ENTERED
    13
                  (A) ADDRESSEE: Human Genome Sciences, Inc.
    14
                  (B) STREET: 9410 Key West Avenue
    15
                  (C) CITY: Rockville
    16
                  (D) STATE: MD
    17
                  (E) COUNTRY: US
    18
                  (F) ZIP: 20850
    19
             (V) COMPUTER READABLE FORM:
    21
                  (A) MEDIUM TYPE: Floppy disk
    22
                  (B) COMPUTER: IBM PC compatible
     23
                  (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     24
                  (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
     25
            (vi) CURRENT APPLICATION DATA:
     27
                  (A) APPLICATION NUMBER: US/09/923,444A
C--> 28
                  (B) FILING DATE: 08-Aug-2001
C-->29
                  (C) CLASSIFICATION:
     30
           (vii) PRIOR APPLICATION DATA:
     32
                  (A) APPLICATION NUMBER: 09/062,815
     33
                  (B) FILING DATE: 199-12-20
     34
          (viii) ATTORNEY/AGENT INFORMATION:
     36
                  (A) NAME: Michele M. Wales
     37
                  (B) REGISTRATION NUMBER: 43,975
     38
                  (C) REFERENCE/DOCKET NUMBER: PF116
     39
            (ix) TELECOMMUNICATION INFORMATION:
     41
                   (A) TELEPHONE: (301) 309-8504
     42
                   (B) TELEFAX: (301) 309-8439
     43
        (2) INFORMATION FOR SEQ ID NO: 1:
     46
             (i) SEQUENCE CHARACTERISTICS:
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                   (A) LENGTH: 2485 base pairs
     49
                   (B) TYPE: nucleic acid
     50
                   (C) STRANDEDNESS: single
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                   (D) TOPOLOGY: linear
     52
             (ii) MOLECULE TYPE: DNA (genomic)
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             (ix) FEATURE:
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                   (A) NAME/KEY: CDS
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                   (B) LOCATION: 266..2446
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             (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/923,444A

DATE: 02/06/2002 TIME: 08:56:41

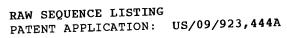
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68 CCAGCCCCAG CCGCCGGAGT GCACGGGCCG AGTCTGCGCG ACTACCCACG CGTGACAGGT 68 CGCAGCTGCT CGGCCGGAGT GCACGGGCCG AGTCTGCGCG ACTACCCACG CGTGACAGGT 70 CCCTGAATGA GAAGGAGCTG ACAGCAGCTG AATTCCATCT TCTCTGTGTG CTGGGGAGCA	240
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81 Ala Leu Glu Glu Pro Val Asp Tyr Lys Gln Ser Val Leu Asn Val Ala	
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85 Gly Glu Ala Gly Gly Lys Gln Lys Ala Vai Glu Glu Glu Leu Asp Ala	
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96 TGC CAG AAA AAT GGA GGA GGT GCT TAC CTG GTG CCC TAC CTG GTG CTG	500
97 Cys Gln Lys Asn Gly Gly Gly Ala Tyr Leu Val Pro Tyr Leu Val 105	
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110 CCC CEC TATE TATE AND CTC ATC ATC GGG TGG AGC ATC TTC TAT TTC TTC	772
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114 155 160	
116 ANG THE CAG TAC CCG CTG CCC TGG AGT GAA TGT CCT GTC GTC AGG	820
117 Lys Ser Phe Gln Tyr Pro Leu Pro Trp Ser Glu Cys Pio Vai Vai Arg	
118 170 175	0.50
120 AND CCC ACC GTC GCA GTG GTG GAG GCA GAG TGT GAA AAG AGC TCA GCC	868
121 Asn Gly Ser Val Ala Val Val Glu Ala Glu Cys Glu Lys Ser Ser Alu	
122 190 195	016
124 ACM ACC MAC MMC MMC MGG MAC CGA GAG GCT TTG GAC ATC TCT GAC TCC ATC	916
125 Thr Thr Tyr Phe Trp Tyr Arg Glu Ala Leu Asp lie Sei Asp Sei lie	
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128 MCC CAC AGE GGC CTC AAC TGG AAG ATG ACC CTG TGC CTC GTG	JU4
129 Ser Glu Ser Gly Gly Leu Asn Trp Lys Met Thr Leu Cys Leu Leu Val	
130 220 225 230	



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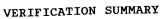
DATE: 02/06/2002 TIME: 08:56:41

· 197 Lys Val Pro Lys Glu Met Phe Thr Val Gly Cys Cys Val Phe Thr Phe											
198 490 THE THE CHE CAG CGC TCC GGA AAC TAC TTT GTC ACC 182	3										
200 CTC GTG GGA CTG TTG TIC GIC CAS GOOD TO THE VAL THE 201 Leu Val Gly Leu Leu Phe Val Gln Arg Ser Gly Asn Tyr Phe Val Thr 520											
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202 TIG TIG TO COC ACC CTG CCA CTC ACT CTC ATC TO	6.										
204 ATG TTC GAT GAC TCA GCC ACG CTO CON THE Leu Ile Val Ile 205 Met Phe Asp Asp Tyr Ser Ala Thr Leu Pro Leu Thr Leu Ile Val Ile 530 535											
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200 THE CONTROL COC TICK ATT TAT GGA CCC AAG AAG TIC AIG	4										
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221 Ala Ser Ile Ile Gln Leu Gly Val Thi Fio Fio Ala 17-											
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222 590 21 224 ATC AAG GAG GAG GCT GCC GAG CGC TAC CTG TAT TTC CCC AAC TGG CCC 21											
225 Tie Lys Glu Glu Ala Ala Glu Arg Tyr Deu Tyr Tho											
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226 605 CTC ATC GTG GTG GCG ACG CTG CCC ATC CCT 21 228 ATG GCA CTC CTG ATC ACC CTC ATC GTG GCG ACG CTG CCC ATC CCT 21											
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232 GTG GTG TTC GTC CTG CGG CAC TTC CAC GTG GTG GTG STC ASP GTG GTG GTG GTG GTG GTG GTG GTG GTG GT											
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238 630 CAR CAR ACC CGC TTC ATC CTC AGC AAG GTG CCC 25	80										
240 AAC CTG GAG GAG AAC GAT GAG ACC CGG 175 HILL Leu Ser Lys Val Pro 241 Asn Leu Glu Glu Asn Asp Glu Thr Arg Phe Ile Leu Ser Lys Val Pro 675 680											
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242 THE THE GOLD ATT. COC ACT CAC CGT TCC TAT CTG GGG CCC	, , ,										
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249 Gly Ser Thr Ser Pro Leu Glu Thi Ser 119 hon 120											
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THE THE COURT COURT ACC COT GAG TOT GAG CIG											
253 Gly Arg Gly Tyr Leu Leu Ala Ser in Fio Gla 305											
254 715 /20	485										
256 TGACCACTGC CCAAGCCCAT GCCCGCTCTC CCCCCACCG											
259 (2) INFORMATION FOR SEQ 1D NO: 2:											
261 (i) SEQUENCE CHARACTERISTICS:											
262 (A) LENGTH: 727 amino acids											
263 (B) TYPE: amino acid											

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/923,444A

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304 306 Gl		<i>C</i> 1	TOO	Glu	Lvs	Ser	Ser	Ala	Thr	Thr	Tyr	Phe	Trp	Tyr	Arg
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307 309 Gl	21-	195	7 cm	Tlo	Ser	Asp	Ser	Ile	ser	Glu	Ser	Gly	Gly	Leu	Asn
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PATENT APPLICATION: US/09/923,444A

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L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]